# **GENOTYPING BY PCR PROTOCOL** MUTANT MOUSE REGIONAL RESOURCE CENTER: UC DAVIS

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530-754-MMRRC

# NAME OF PCR: STOCK Fgf4<sup>tm1.2Mrt</sup>/Mmucd

## **MMRRC: 011204-UCD**

### **Protocol:**

Reagent/Constituent	Volume (µL)
Water	11.275
10x Buffer	2.5
MgCl <sub>2</sub> (stock concentration is 25mM)	1.7
Betaine (stock concentration is 5M) Optional	6.5
dNTPs (stock concentration is 10mM)	0.5
DMSO Optional	0.325
Primer 1. (stock concentration is 20µM)	0.5
Primer 2. (stock concentration is 20µM)	0.5
Taq Polymerase 5Units/µL	0.2
DNA (example) extracted w/ "Qiagen DNeasy columns or other similar silica based kits"	1.0
TOTAL VOLUME OF REACTION:	25.000 μL

### Comments on protocol:

- Protocol may work with other DNA extraction methods. •
- Use Touch-Down cycling protocol-first 10 cycles anneal at 65°C decreasing in temperature by 1.0°C; next 30 cycles anneal at 55°C.
- Betaine and DMSO have been standardized due to high GC content. Protocol may be tested without. Also, may adjust .

#### Strategy:

Steps		Temp (°C )	Time (m:ss)	# of Cycles
1. Initiation/Melting	HOT START?	94	5:00	1
2. Denaturation		94	0:15	
3. Annealing	steps 2-3-4 cycle in sequence	65 to 55 (↓1°C/cycle)	0:30	40x
4. Elongation		72	0:40	
5. Amplification		72	5:00	1
6. Finish		15	$\infty$	n/a

### **Primers:**

imers:		Electrophoresis Protocol:			
Name	Nucleotide Sequence (5' - 3')	Argarose: 1.5%	V: 90		
1. EM26	TCTGGAGAGGAACTAGGAATGG	Estimated Running:Tin	Estimated Running:Time: 90 min.		
2. EM27	GAAGAGAAGCAGGCAGATGC	Primer Combination	Band (bp)	Genotype	
		1 and 2	344	WT	

1 and 2

396

Floxed

